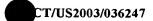


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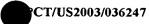
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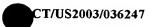
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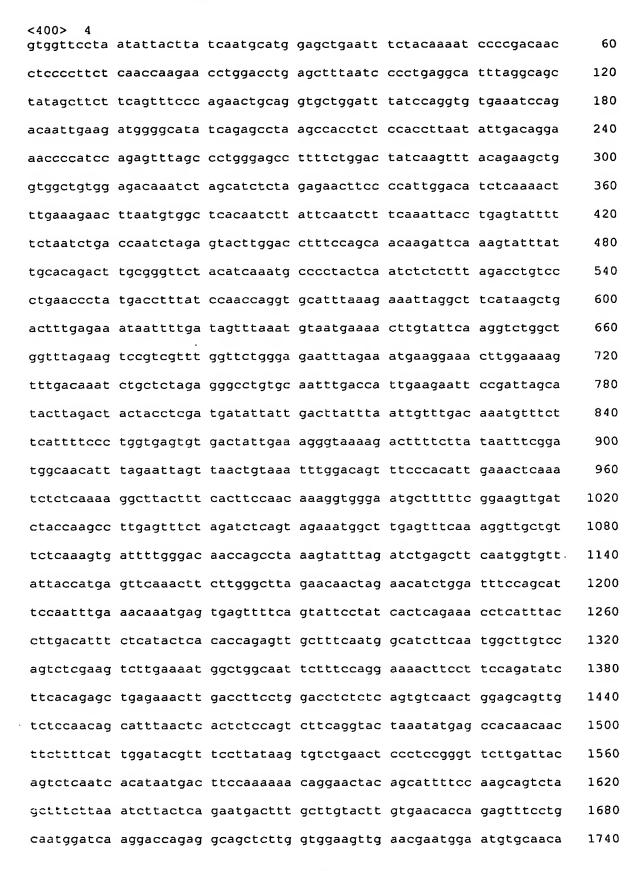
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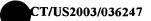
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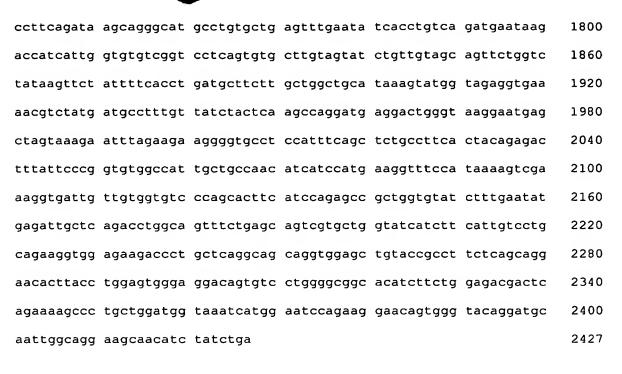
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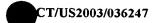
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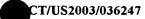
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Phe Asp Lys Ser Ala Leu Glu Gly Leu Cys Asn Leu Thr Ile Glu Glu 245 250 255

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Ser Leu Lys Arg Leu Thr Phe Thr Ser Asn Lys Gly Gly Asn Ala Phe 325 330 335

Ser Glu Val Asp Leu Pro Ser Leu Glu Phe Leu Asp Leu Ser Arg Asn 340 345 350

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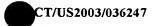
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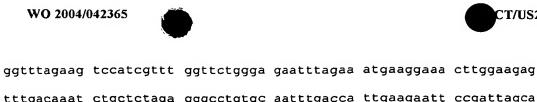
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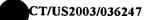
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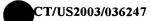


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agt gtc ctg ggg cgg cac att ttc tgg aga cga ctc aga aaa gcc ctg Ser Val Leu Gly Arg His Ile Phe Trp Arg Arg Leu Arg Lys Ala Leu 770 775 . 780	352
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- Asn Pro Leu Arg His Leu Gly Ser Tyr Ser Phe Phe Ser Phe Pro Glu 35 40 45
- Leu Gln Val Leu Asp Leu Ser Arg Cys Glu Ile Gln Thr Ile Glu Asp 50 55 60
- Gly Ala Tyr Gln Ser Leu Ser Leu Ser Thr Leu Ile Leu Thr Gly 70 75 80
- Asn Pro Ile Gln Ser Leu Ala Leu Gly Ala Phe Ser Gly Leu Ser Ser 85 90 95
- Leu Gln Lys Leu Val Ala Val Glu Thr Asn Leu Ala Ser Leu Glu Asn 100 105 110
- Phe Pro Ile Gly His Leu Lys Thr Leu Lys Glu Leu Asn Val Ala His 115 120 125
- Asn Leu Ile Gln Ser Phe Lys Leu Pro Glu Tyr Phe Ser Asn Leu Thr 130 135 140
- Asn Leu Glu His Leu Asp Leu Ser Ser Asn Lys Ile Gln Ser Ile Tyr 145 150 155 160
- Cys Lys Asp Leu Gln Val Leu His Gln Met Pro Leu Leu Asn Leu Ser 165 170 175
- Leu Asp Leu Ser Leu Asn Pro Met Asn Phe Ile Gln Pro Gly Ala Phe 180 185 190
- Lys Glu Ile Ser Leu Xaa Lys Leu Thr Leu Arg Asn Asn Phe Asp Ser 195 200 205
- Leu Asn Val Met Lys Thr Cys Ile Gln Gly Leu Ala Gly Leu Glu Val 210 215 220
- His Arg Leu Val Leu Gly Glu Phe Arg Asn Glu Gly Asn Leu Glu Glu 225 230 235 240
- Phe Asp Lys Ser Ala Leu Glu Gly Leu Cys Asn Leu Thr Ile Glu Glu 245 250 255



Phe Arg Leu Ala Tyr Leu Asp His Tyr Leu Asp Asp Ile Ile Asp Leu 260 265 270

Phe Asn Cys Leu Ala Asn Val Ser Ser Phe Ser Leu Val Ser Val Thr 275 280 285

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Glu Leu Val Asn Cys Lys Phe Gly Gln Phe Pro Thr Leu Asn Leu Lys 305 310 315 320

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Ser Glu Val Asp Leu Pro Ser Leu Glu Phe Leu Asp Leu Ser Arg Asn 340 345 350

Gly Leu Ser Phe Lys Gly Cys Cys Ser Gln Ser Asp Phe Gly Thr Asn 355 360 365

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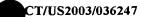
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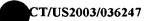
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- Lys Lys Gln Glu Leu Gln Arg Phe Pro Ser Ser Leu Ala Phe Leu Asn 530 535 540
- Leu Thr Gln Asn Asp Phe Ala Cys Thr Cys Glu His Glu Ser Phe Leu 545 550 555 560
- Gln Trp Ile Lys Asp Gln Arg Gln Leu Leu Val Glu Val Glu Arg Met 565 570 575
- Glu Cys Ala Thr Pro Ser Asp Lys Gln Gly Met Pro Val Leu Ser Leu 580 585 590
- Asn Ile Thr Cys Gln Met Asn Lys Thr Ile Ile Gly Val Ser Val Leu 595 600 605
- Ser Val Leu Val Val Ser Val Val Ala Val Leu Val Tyr Lys Phe Tyr 610 615 620
- Phe His Leu Met Leu Leu Ala Gly Cys Met Lys Tyr Gly Arg Gly Glu 625 630 640
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- Val Arg Asn Glu Leu Val Lys Asn Leu Glu Glu Gly Val Pro Pro Phe 660 665 670
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- Ala Asn Ile Ile His Glu Gly Phe His Lys Ser Arg Lys Val Ile Val 690 700
- Val Val Ser Gln His Phe Ile Gln Ser Arg Trp Cys Ile Phe Glu Tyr 705 710 715 720
- Fig 11e Ala Gln Thr Trp Gln Phe Leu Ser Ser His Ala Gly Ile Ile 725 730 735



Phe Ile Val Leu Gln Lys Val Glu Lys Thr Leu Leu Arg Gln Gln Val 740 745 750

Glu Leu Tyr Arg Leu Leu Ser Arg Asn Thr Tyr Leu Glu Trp Glu Asp 755 760 . 765

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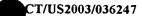
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ggc tt Gly Le					-	_			-	_					1104
agc ct Ser Le 37	u Lys														1152
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aaa aa Lys As 465															1440



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		aag Lys								172	28
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		tgt Cys								183	24
		gtg Val								18	72
		atg Met								19	20
		gat Asp								19	68
		gaa Glu 660								20	16
		ctt Leu								20	64
	Ile	atc Ile					Lys			21	12
		cag Gln		Ile						21	60



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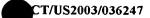




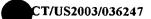
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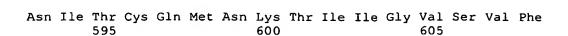
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- Lys Asn Leu Thr Phe Leu Asp Leu Ser Gln Cys Gln Leu Glu Gln Leu 465 470 475 480
- Ser Pro Thr Ala Phe Asp Thr Leu Asn Lys Leu Gln Val Leu Asn Met 485 490 495
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- Leu Thr Gln Asn Asp Phe Ala Cys Thr Cys Glu His Gln Ser Phe Leu 545 550 555 560
- Gln Trp Ile Lys Asp Gln Arg Gln Leu Leu Val Glu Ala Glu Arg Met 565 570 575
- Glu Cys Ala Thr Pro Ser Asp Lys Gln Gly Met Pro Val Leu Ser Leu 580 585 590





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Phe His Leu Met Leu Leu Ala Gly Cys Ile Xaa Tyr Gly Arg Gly Glu 625 635 640

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Gln Leu Cys Leu His Tyr Arg Asp Phe Ile Pro Gly Val Ala Ile Ala 675 680 685

Ala Asn Ile Ile His Glu Gly Phe His Lys Ser Arg Lys Val Ile Val 690 695 700

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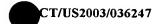
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336

384

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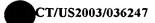
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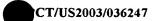
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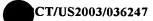
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- Asp Thr Phe Pro Tyr Lys His Leu Tyr Ser Leu His Val Leu Asp Tyr 450 455 460
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Phe Ile Pro Gly Val Ala Ile Ala Ala Asn Ile Ile His Glu Gly Phe 630

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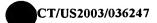
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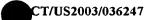
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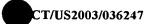
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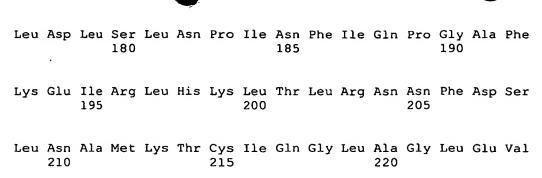
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Phe Arg Leu Ala Tyr Leu Asp Asp Phe Leu Asp Asp Ile Ile Asp Leu 260 265 270

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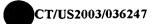
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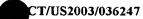
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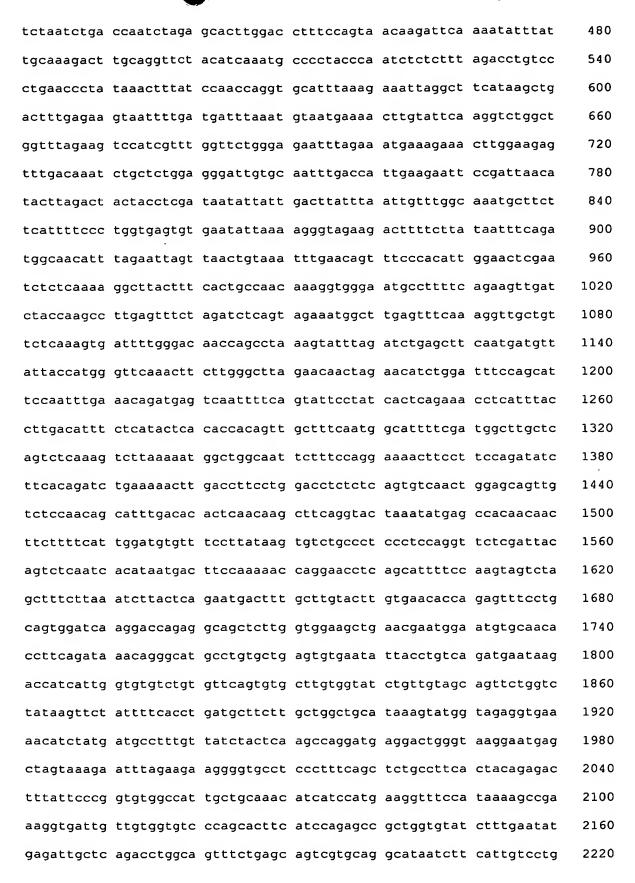
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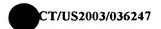
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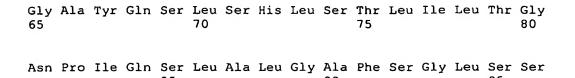


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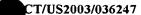
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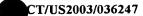
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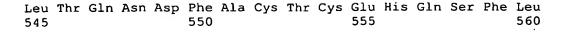




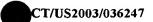
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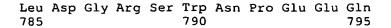
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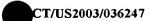
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Leu Gln Val Leu Asp Leu Ser Arg Cys Glu Ile Gln Thr Ile Glu Asp





50 55 60 ggg gca tat cag agc cta agc cac ctc tcc acc tta ata ttg aca gga 240 Gly Ala Tyr Gln Ser Leu Ser His Leu Ser Thr Leu Ile Leu Thr Gly 75 288 aac ccc atc cag agt tta gcc ctg gga gcc ttt tct gga cta tca agt Asn Pro Ile Gln Ser Leu Ala Leu Gly Ala Phe Ser Gly Leu Ser Ser 336 tta cag aag ctg gtg gct gtg gag aca aat cta gca tct cta gag aac Leu Gln Lys Leu Val Ala Val Glu Thr Asn Leu Ala Ser Leu Glu Asn 100 105 ttc ccc att gga cat ctc aaa act ttg aaa gaa ctt aat gtg gct cac 384 Phe Pro Ile Gly His Leu Lys Thr Leu Lys Glu Leu Asn Val Ala His 120 aat ctt atc caa tct ttc aaa tta cct gag tat ttt tct aat ctg acc 432 Asn Leu Ile Gln Ser Phe Lys Leu Pro Glu Tyr Phe Ser Asn Leu Thr 135 aat cta gag cac ttg gac ctt tcc agc aac aag att caa agt att tat 480 Asn Leu Glu His Leu Asp Leu Ser Ser Asn Lys Ile Gln Ser Ile Tyr 145 tgc aca gac ttg cgg gtt cta cat caa atg ccc cta ctc aat ctc tct 528 Cys Thr Asp Leu Arg Val Leu His Gln Met Pro Leu Leu Asn Leu Ser 170 tta gac ctg tcc ctg aac cct atg aac ttt atc caa cca ggt gca ttt 576 Leu Asp Leu Ser Leu Asn Pro Met Asn Phe Ile Gln Pro Gly Ala Phe 180 185 aaa gaa att agg ctt cat aag ctg act ttg aga aat aat ttt gat agt 624 Lys Glu Ile Arg Leu His Lys Leu Thr Leu Arg Asn Asn Phe Asp Ser 195 200 tta aat gta atg aaa act tgt att caa ggt ctg gct ggt tta gaa gtc 672 Leu Asn Val Met Lys Thr Cys Ile Gln Gly Leu Ala Gly Leu Glu Val 210 cat cqt ttq qtt ctq qqa qaa ttt aga aat qaa qaa aac ttq qaa aag 720 His Arg Leu Val Leu Gly Glu Phe Arg Asn Glu Glu Asn Leu Glu Lys 225 230 240 ttt gac aaa tct gct cta gag ggc ctg tgc aat ttg acc att gaa gaa 768 Phe Asp Lys Ser Ala Leu Glu Gly Leu Cys Asn Leu Thr Ile Glu Glu ttc cga tta gca tac tta gac tac tac ctc gat gat att att gac tta 816 Phe Arg Leu Ala Tyr Leu Asp Tyr Tyr Leu Asp Asp Ile Ile Asp Leu ttt aat tgt ttg aca aat gtt tct tca ttt tcc ctg gtg agt gtg act 864 Phe Asn Cys Leu Thr Asn Val Ser Ser Phe Ser Leu Val Ser Val Thr 275 280 att aaa agc gta aaa gac ttt tct tat aat ttc gga tgg caa cat tta 912 Ile Lys Ser Val Lys Asp Phe Ser Tyr Asn Phe Gly Trp Gln His Leu



	290					295					300					
						ttt Phe										960
tct Ser	ctc Leu	aaa Lys	agg Arg	ctt Leu 325	act Thr	ttc Phe	act Thr	tcc Ser	aac Asn 330	aaa Lys	ggt Gly	GJ À āāā	aat Asn	gct Ala 335	ttt Phe	1008
						agc Ser										1056
ggc Gly	ttg Leu	agt Ser 355	ttc Phe	aaa Lys	ggt Gly	tgc Cys	tgt Cys 360	tct Ser	caa Gln	agt Ser	gat Asp	ttt Phe 365	ggg Gly	aca Thr	acc Thr	1104
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						agt Ser										1248
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						tca Ser										1488
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												agt Ser		1	.680
												gaa Glu		1	.728
												ctg Leu 590		1	.776
		-	_	-		_						tcg Ser	-	1	824
-		-	-			-	_	_	-	-		aag Lys		1	.872
												aga Arg		1	.920
												gag Glu		1	.968
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_				_					_	_	_	gtg Val		 2	2112
												ttt Phe		2	2160
												ggt Gly		2	208
												cgg Arg 750		2	256
												tgg Trp		2	2304
												aaa Lys		2	2352

770 775

ctg gat ggt aaa tca tgg aat cca gaa gga aca gtg ggt aca gga tgc 2400 Leu Asp Gly Lys Ser Trp Asn Pro Glu Gly Thr Val Gly Thr Gly Cys 785 790 795 800

780

aat tgg cag gaa gca aca tct atc tga 2427 Asn Trp Gln Glu Ala Thr Ser Ile 805

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<213> Pan troglodytes

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<222> (758)..(758)

<223> The 'Xaa' at location 758 stands for Leu.

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Ile Pro Asp Asn Leu Pro Phe Ser Thr Lys Asn Leu Asp Leu Ser Phe
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Asn Pro Leu Arg His Leu Gly Ser Tyr Ser Phe Phe Ser Phe Pro Glu 35 40 45

Leu Gln Val Leu Asp Leu Ser Arg Cys Glu Ile Gln Thr Ile Glu Asp 50 55 60

Gly Ala Tyr Gln Ser Leu Ser His Leu Ser Thr Leu Ile Leu Thr Gly 65 70 75 80

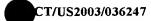
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Leu Gln Lys Leu Val Ala Val Glu Thr Asn Leu Ala Ser Leu Glu Asn 100 105 110

Phe Pro Ile Gly His Leu Lys Thr Leu Lys Glu Leu Asn Val Ala His 115 120 125

Asn Leu Ile Gln Ser Phe Lys Leu Pro Glu Tyr Phe Ser Asn Leu Thr 130 135 140

Asn Leu Glu His Leu Asp Leu Ser Ser Asn Lys Ile Gln Ser Ile Tyr



145 150 155 160

Cys Thr Asp Leu Arg Val Leu His Gln Met Pro Leu Leu Asn Leu Ser 165 170 175

Leu Asp Leu Ser Leu Asn Pro Met Asn Phe Ile Gln Pro Gly Ala Phe 180 185 190

Lys Glu Ile Arg Leu His Lys Leu Thr Leu Arg Asn Asn Phe Asp Ser 195 200 205

Leu Asn Val Met Lys Thr Cys Ile Gln Gly Leu Ala Gly Leu Glu Val 210 215 220

His Arg Leu Val Leu Gly Glu Phe Arg Asn Glu Glu Asn Leu Glu Lys 225 230 235 240

Phe Asp Lys Ser Ala Leu Glu Gly Leu Cys Asn Leu Thr Ile Glu Glu 245 250 255

Phe Arg Leu Ala Tyr Leu Asp Tyr Tyr Leu Asp Asp Ile Ile Asp Leu 260 265 270

Phe Asn Cys Leu Thr Asn Val Ser Ser Phe Ser Leu Val Ser Val Thr 275 280 285

Ile Lys Ser Val Lys Asp Phe Ser Tyr Asn Phe Gly Trp Gln His Leu 290 295 300

Glu Leu Val Lys Cys Lys Phe Gly Gln Phe Pro Thr Leu Lys Leu Lys 305 310 315 320

Ser Leu Lys Arg Leu Thr Phe Thr Ser Asn Lys Gly Gly Asn Ala Phe 325 330 335

Ser Glu Val Asp Leu Pro Ser Leu Glu Phe Leu Asp Leu Ser Arg Asn 340 345 350

Gly Leu Ser Phe Lys Gly Cys Cys Ser Gln Ser Asp Phe Gly Thr Thr 355 360 365

Ser Leu Lys Tyr Leu Asp Leu Ser Phe Asn Gly Val Ile Thr Met Ser 370 380

Ser Asn Phe Leu Gly Leu Glu Gln Leu Glu His Leu Asp Phe Gln His

385 390 395 400

Ser Asn Leu Lys Gln Met Ser Glu Phe Ser Val Phe Leu Ser Leu Arg 405 410 415

Asn Leu Ile Tyr Leu Asp Ile Ser His Thr His Thr Arg Val Ala Phe 420 425 430

Asn Gly Ile Phe Asn Gly Leu Ser Ser Leu Glu Val Leu Lys Met Ala 435 440 445

Gly Asn Ser Phe Gln Glu Asn Phe Leu Pro Asp Ile Phe Thr Glu Leu 450 460

Arg Asn Leu Thr Phe Leu Asp Leu Ser Gln Cys Gln Leu Glu Gln Leu 465 470 475 480

Ser Pro Thr Ala Phe Asn Ser Leu Ser Ser Leu Gln Val Leu Asn Met 485 490 495

Ser His Asn Asn Phe Phe Ser Leu Asp Thr Phe Pro Tyr Lys Cys Leu 500 505 510

Asn Ser Leu Gln Val Leu Asp Tyr Ser Leu Asn His Ile Met Thr Ser 515 520 525

Lys Lys Gln Glu Leu Gln His Phe Pro Ser Ser Leu Ala Phe Leu Asn 530 540

Leu Thr Gln Asn Asp Phe Ala Cys Thr Cys Glu His Gln Ser Phe Leu 545 550 555 560

Gln Trp Ile Lys Asp Gln Arg Gln Leu Leu Val Glu Val Glu Arg Met 565 570 575

Glu Cys Ala Thr Pro Ser Asp Lys Gln Gly Met Pro Val Leu Ser Leu 580 585 590

Asn Ile Thr Cys Gln Met Asn Lys Thr Ile Ile Gly Val Ser Val Leu 595 600 605

Ser Val Leu Val Val Ser Val Val Ala Val Leu Val Tyr Lys Phe Tyr 610 615 620

Fhe His Leu Met Leu Leu Ala Gly Cys Ile Lys Tyr Gly Arg Gly Glu

625 630 635 640

Asn Ile Tyr Asp Ala Phe Val Ile Tyr Ser Ser Gln Asp Glu Asp Trp 645 650 655

Val Arg Asn Glu Leu Val Lys Asn Leu Glu Glu Gly Val Pro Pro Phe 660 665 670

Gln Leu Cys Leu His Tyr Arg Asp Phe Ile Pro Gly Val Ala Ile Ala 675 680 685

Ala Asn Ile Ile His Glu Gly Phe His Lys Ser Arg Lys Val Ile Val 690 695 700

Val Val Ser Gln His Phe Ile Gln Ser Arg Trp Cys Ile Phe Glu Tyr 705 710 715 720

Glu Ile Ala Gln Thr Trp Gln Phe Leu Ser Ser Arg Ala Gly Ile Ile 725 730 735

Phe Ile Val Leu Gln Lys Val Glu Lys Thr Leu Leu Arg Arg Gln Val 740 745 750

Glu Leu Tyr Arg Leu Xaa Ser Arg Asn Thr Tyr Leu Glu Trp Glu Asp 755 760 765

Ser Val Leu Gly Arg His Ile Phe Trp Arg Arg Leu Arg Lys Ala Leu 770 775 780

Leu Asp Gly Lys Ser Trp Asn Pro Glu Gly Thr Val Gly Thr Gly Cys 785 790 795 800

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<213> Amino acid

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Leu Asp Asp Ile Ile Asp Leu Phe Asn Cys Leu Ala Asn Ala Ser Ser 20 25 30

Phe Ser Leu 35

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Phe Ser Leu 35

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Phe Ser Leu

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Leu Asp Asp Ile Ile Asp Leu Phe Asn Cys Leu Ala Asn Ala Ser Ser 20 25 30

Phe Ser Leu 35

<210> 30

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Phe Ser Leu 35

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Phe Ser Leu 35 <210> 32

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Leu Asp Asn Ile Ile Asp Leu Phe Asn Cys Leu Ala Asn Ala Ser Ser 25

Phe Ser Leu 35

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Leu Asp Asn Ile Ile Asp Leu Phe Asn Cys Leu Ala Asn Ala Ser Ser

Phe Ser Leu 35

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Ser Ser Lys Asp Ser Ile Asp Leu Leu Asn Cys Leu Ala Asp Ile Ser

Lys Ile Ser Leu 35

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WO 2004/042365



<213> Amino acid

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Ser Gly Asp Asp Thr Asp Leu Phe Asn Cys Leu Ala Asn Val Ser Val

Ile Ser Leu 35

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Ile Ser Leu 35

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Ile Ser Leu 35

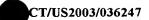
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Ser Leu